

1/4

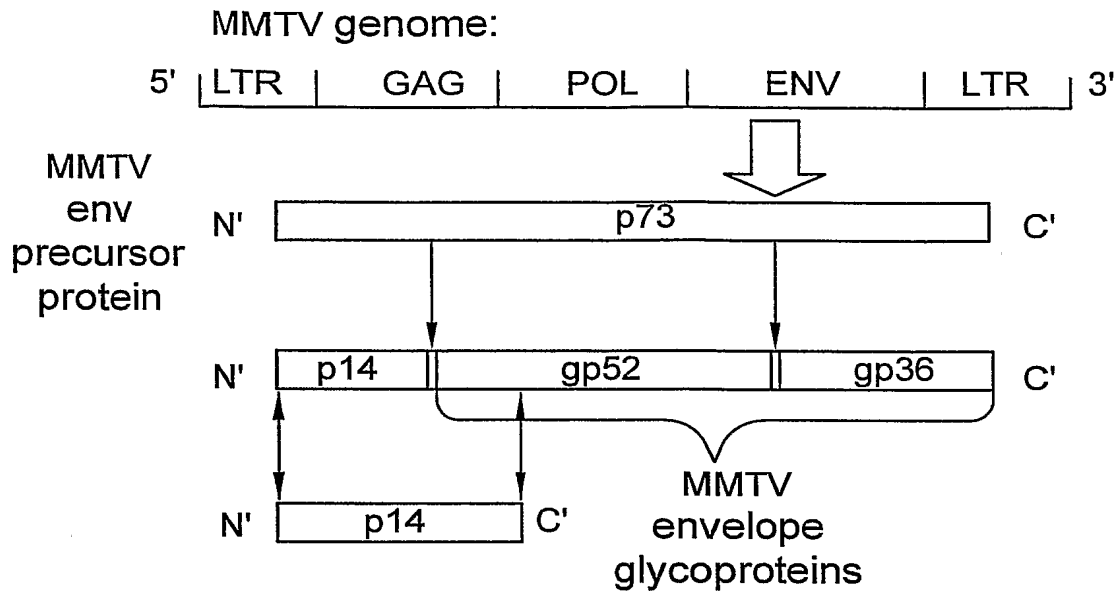


FIG. 1A

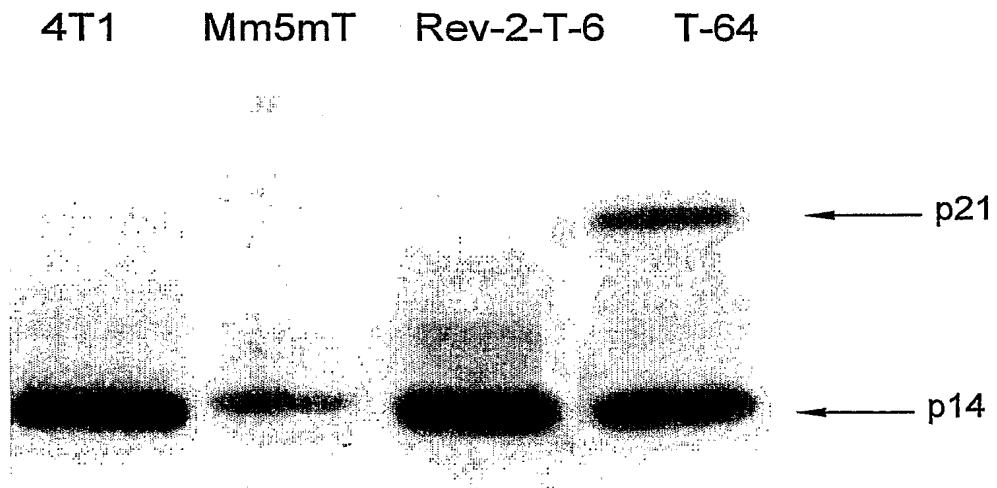


FIG. 1B

af248270 2 GSLTDSSDLSGKDRRPRLALRRKRRREMRKITKKVRRMTLDPIKEKA 51  
||| ||| ||| ||| . ||| ||| ||| ||| : ||| ||| ||| ||| |||  
p14 7 GSPTGSSDLLSGKKQORPHLALRRKRRREMKKINRKVRRMNLDLIKEKA 56  
52 WQQLQALISEAEEVLKTSQTPQTSLTLEFLALLSVLGPLPVTG 93  
||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||  
57 WQHLQALISEAEEVLKTSQTPQTSLTLEFLALLSVLGPPPVTG 98

**FIG. 2**



FIG.3A

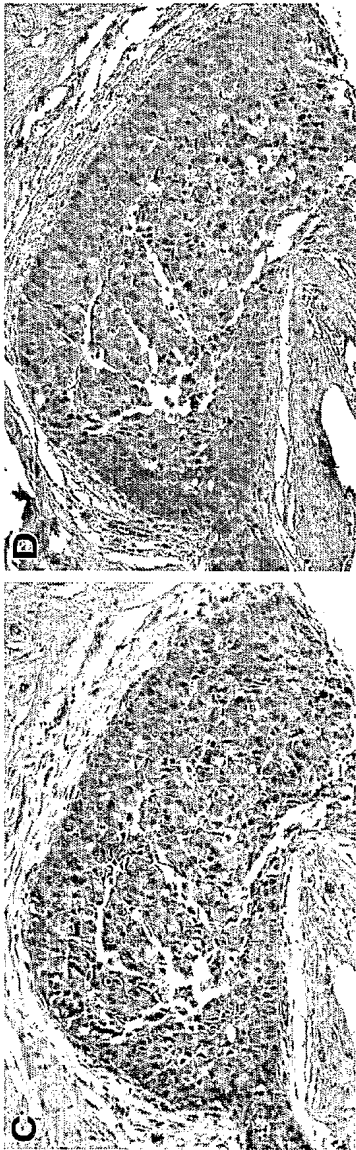


FIG.3C

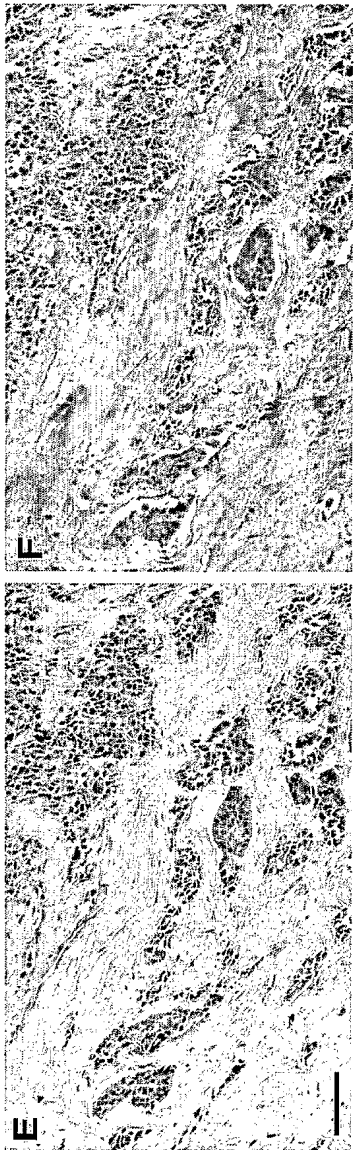


FIG.3E

FIG.3B

FIG.3D

FIG.3F

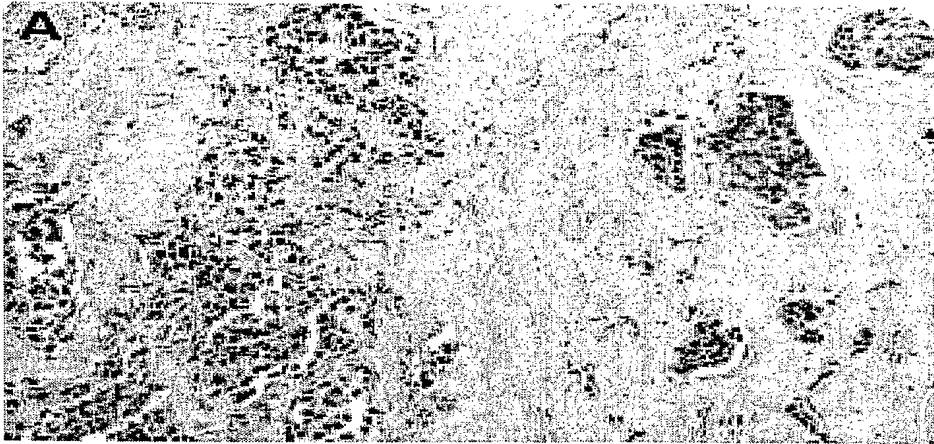


FIG. 4A

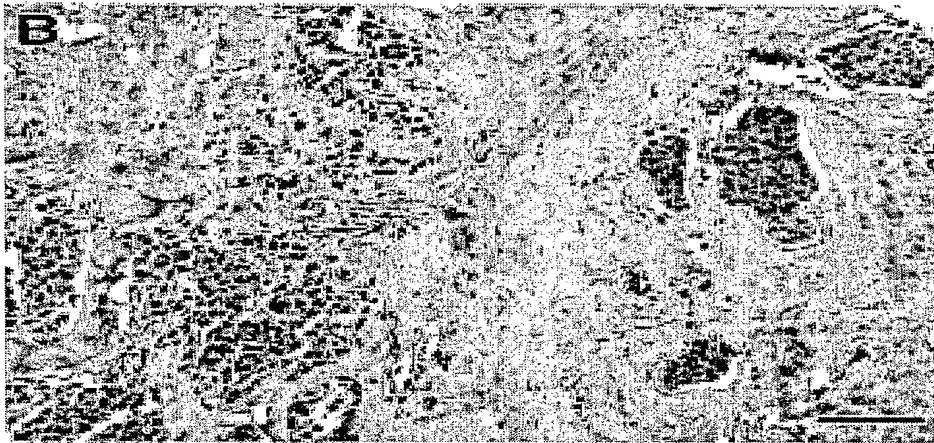


FIG. 4B



FIG. 4C